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SEQUENCE LISTING

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<120> RECEPTOR REGULATOR

<130> 3116USOP

<150> PCT/JP03/14101

<151> 2003-11-05

<150> JP 2002-322534

<151> 2002-11-06

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Gln Thr Glu Leu Phe Met Pro Ile Cys Ala Thr Tyr Leu Leu Ile Phe
           35           40           45
Val Val Gly Ala Val Gly Asn Gly Leu Thr Cys Leu Val Ile Leu Arg
           50           55           60
His Lys Ala Met Arg Thr Pro Thr Asn Tyr Tyr Leu Phe Ser Leu Ala
65           70           75           80
Val Ser Asp Leu Leu Val Leu Leu Val Gly Leu Pro Leu Glu Leu Tyr
           85           90           95
Glu Met Trp His Asn Tyr Pro Phe Leu Leu Gly Val Gly Gly Cys Tyr
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Phe Arg Thr Leu Leu Phe Glu Met Val Cys Leu Ala Ser Val Leu Asn
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Val Thr Ala Leu Ser Val Glu Arg Tyr Val Ala Val Val His Pro Leu
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Gln Ala Arg Ser Met Val Thr Arg Ala His Val Arg Arg Val Leu Gly
145          150          155          160
Ala Val Trp Gly Leu Ala Met Leu Cys Ser Leu Pro Asn Thr Ser Leu
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His Gly Ile Arg Gln Leu His Val Pro Cys Arg Gly Pro Val Pro Asp
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Ser Ala Val Cys Met Leu Val Arg Pro Arg Ala Leu Tyr Asn Met Val
           195          200          205
Val Gln Thr Thr Ala Leu Leu Phe Phe Cys Leu Pro Met Ala Ile Met
           210          215          220
Ser Val Leu Tyr Leu Leu Ile Gly Leu Arg Leu Arg Arg Glu Arg Leu
225          230          235          240
Leu Leu Met Gln Glu Ala Lys Gly Arg Gly Ser Ala Ala Ala Arg Ser

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Ser	Val	Leu	Tyr	Leu	Leu	Ile	Gly	Leu	Arg	Leu	Arg	Arg	Glu	Arg	Met		

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 245 250 255
 Thr Ser His Arg Arg Ile Gln Leu Gln Asp Arg Gly Arg Arg Gln Val
 260 265 270
 Thr Lys Met Leu Phe Ala Leu Val Val Phe Gly Ile Cys Trp Ala
 275 280 285
 Pro Phe His Ala Asp Arg Ile Met Trp Ser Leu Val Tyr Gly His Ser
 290 295 300
 Thr Glu Gly Leu His Leu Ala Tyr Gln Cys Val His Ile Ala Ser Gly
 305 310 315 320
 Ile Phe Phe Tyr Leu Gly Ser Ala Ala Asn Pro Val Leu Tyr Ser Leu
 325 330 335
 Met Ser Thr Arg Phe Arg Glu Thr Phe Leu Gln Ala Leu Gly Leu Gly
 340 345 350
 Thr Gln Cys Cys His Arg Arg Gln Pro Tyr His Gly Ser His Asn His
 355 360 365
 Ile Arg Leu Thr Thr Gly Ser Thr Leu Cys Asp Val Gly His Arg Asn
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 Glu Thr Asp Pro Ser
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 35 40 45
 Ser Val Val Tyr Val Pro Ile Phe Val Val Gly Val Ile Gly Asn Val
 50 55 60
 Leu Val Cys Leu Val Ile Leu Gln His Gln Ala Met Lys Thr Pro Thr
 65 70 75 80
 Asn Tyr Tyr Leu Phe Ser Leu Ala Val Ser Asp Leu Leu Val Leu Leu
 85 90 95
 Leu Gly Met Pro Leu Glu Val Tyr Glu Met Trp Arg Asn Tyr Pro Phe
 100 105 110
 Leu Phe Gly Pro Val Gly Cys Tyr Phe Lys Thr Ala Leu Phe Glu Thr
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 Val Cys Phe Ala Ser Ile Leu Ser Ile Thr Thr Val Ser Val Glu Arg
 130 135 140
 Tyr Val Ala Ile Leu His Pro Phe Arg Ala Lys Leu Gln Ser Thr Arg
 145 150 155 160
 Arg Arg Ala Leu Arg Ile Leu Gly Ile Val Trp Gly Phe Ser Val Leu
 165 170 175
 Phe Ser Leu Pro Asn Thr Ser Ile His Gly Ile Lys Phe His Tyr Phe
 180 185 190
 Pro Asn Gly Ser Leu Val Pro Gly Ser Ala Thr Cys Thr Val Ile Lys
 195 200 205
 Pro Met Trp Ile Tyr Asn Phe Ile Ile Gln Val Thr Ser Phe Leu Phe

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Asn	Ile	Gln	Arg	Pro	Cys	Arg	Lys	Ser	Val	Asn	Lys	Met	Leu	Phe	Val
			260						265					270	
Leu	Val	Leu	Val	Phe	Ala	Ile	Cys	Trp	Ala	Pro	Phe	His	Ile	Asp	Arg
		275						280					285		
Leu	Phe	Phe	Ser	Phe	Val	Glu	Glu	Trp	Ser	Glu	Ser	Leu	Ala	Ala	Val
	290					295						300			
Phe	Asn	Leu	Val	His	Val	Val	Ser	Gly	Val	Phe	Phe	Tyr	Leu	Ser	Ser
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Ala	Val	Asn	Pro	Ile	Ile	Tyr	Asn	Leu	Leu	Ser	Arg	Arg	Phe	Gln	Ala
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Ala	Phe	Gln	Asn	Val	Ile	Ser	Ser	Phe	His	Lys	Gln	Trp	His	Ser	Gln
			340						345				350		
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Cys	His	Phe	Val	Glu	Leu	Thr	Glu	Asp	Ile	Gly	Pro	Gln	Phe	Pro	Cys
	370					375					380				
Gln	Ser	Ser	Met	His	Asn	Ser	His	Leu	Pro	Thr	Ala	Leu	Ser	Ser	Glu
385					390					395					400
Gln	Met	Ser	Arg	Thr	Asn	Tyr	Gln	Ser	Phe	His	Phe	Asn	Lys	Thr	
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